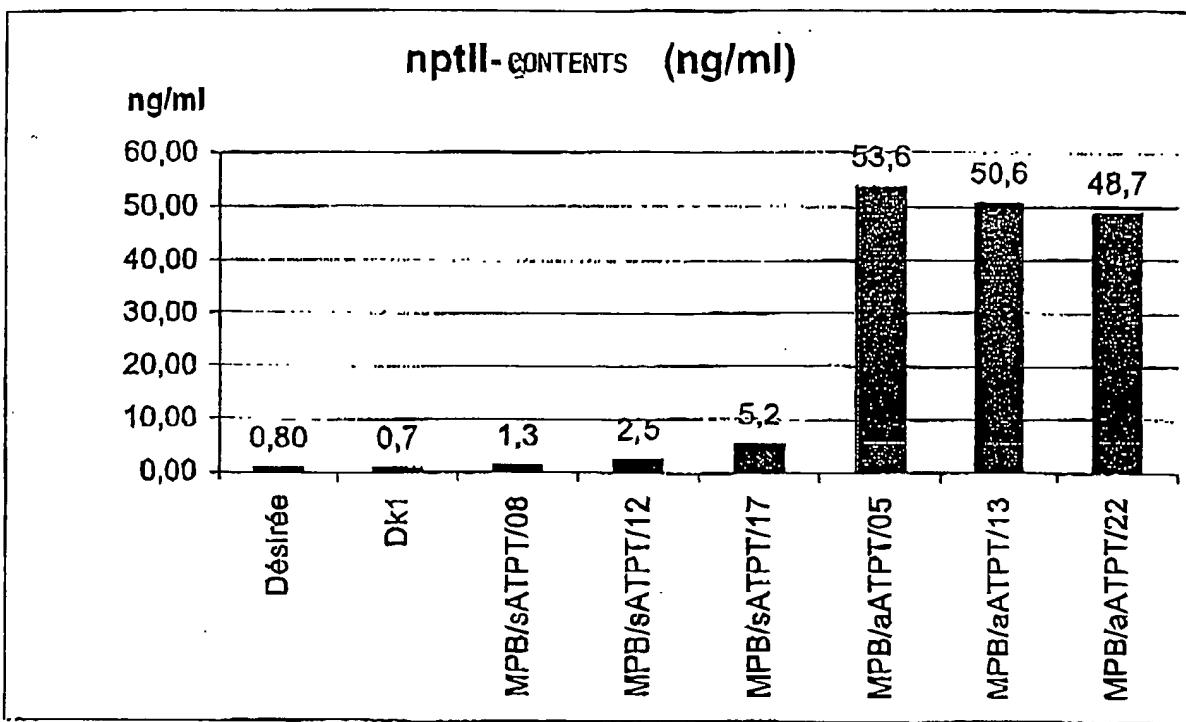


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Figure 1:



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Fig. 2

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atggccccc aacccttct atcttgctt tcacttcct tgcctcttt tacaaggcaca	60
tgtta gca gct gat gtg cag ctg gtg gag tct ggg gga ggc tta gtg Ala Ala Asp Val Gin Leu Val Glu Ser Gly Gly Gly Leu Val 1 5 10	108
cag cct gga ggg tcc cgg aaa ctc tcc tgt gca gcc tct gga ttc act Gln Pro Gly Gly Ser Arg Lys Leu Ser Cys Ala Ala Ser Gly Phe Thr 15 20 25 30	156
ttc agt agc ttt gga atg cac tgg gtt cgt cag gct cca gag aag ggg Phe Ser Ser Phe Gly Met His Trp Val Arg Gln Ala Pro Glu Lys Gly 35 40 45	204
ctg gag tgg gtc gca tat att agt agt ggc agt agt acc atc tac tat Leu Glu Trp Val Ala Tyr Ile Ser Ser Gly Ser Ser Thr Ile Tyr Tyr 50 55 60	252
gca gac aca gtg aag ggc cga ttc acc atc tcc aga gac aat ccc aag Ala Asp Thr Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Pro Lys 65 70 75	300
aac acc ctg ttc ctg caa atg acc agt cta agg tct gag gac acg gcc Asn Thr Leu Phe Leu Gln Met Thr Ser Leu Arg Ser Glu Asp Thr Ala 80 85 90	348
atg tat tac tgc gca aga gat tac ggg gct tat tgg ggc caa ggg acc Met Tyr Tyr Cys Ala Arg Asp Tyr Gly Ala Tyr Trp Gly Gln Gly Thr 95 100 105 110	396
acg gtc acc gtc tcc tca ggt gga ggc ggt tca ggc gga ggt ggc tct Thr Val Thr Val Ser Ser Gly Gly Ser Gly Gly Gly Ser 115 120 125	444
ggc ggt ggc gga tcg gac att gag ctc acc cag tct cca gca atc atg Gly Gly Gly Ser Asp Ile Glu Leu Thr Gln Ser Pro Ala Ile Met 130 135 140	492
tct gca tct cca ggg gag aag gtc acc atg acc tgc agt gcc agt tca Ser Ala Ser Pro Gly Glu Lys Val Thr Met Thr Cys Ser Ala Ser Ser 145 150 155	540
agt gta agg tac atg aac tgg ttc caa cag aag tca ggc acc tcc ccc Ser Val Arg Tyr Met Asn Trp Phe Gln Gln Lys Ser Gly Thr Ser Pro 160 165 170	588
aaa aga tgg att tat gac aca tcc aaa ctg tct tct gga gtc cct gct Lys Arg Trp Ile Tyr Asp Thr Ser Lys Leu Ser Ser Gly Val Pro Ala 175 180 185 190	636
cgc ttc agt ggc agt ggg tct ggg acc tct tac tct ctc aca atc agc Arg Phe Ser Gly Ser Gly Ser Gly Thr Ser Tyr Ser Leu Thr Ile Ser 195 200 205	684
agc atg gag gct gaa gat gct gcc act tat tac tgc cag cag tgg agt Ser Met Glu Ala Glu Asp Ala Ala Thr Tyr Tyr Cys Gln Gln Trp Ser 210 215 220	732
agt aat cca ctc act ttc ggt gct ggg acc aag ctg gag ctg aaa cgg Ser Asn Pro Leu Thr Phe Gly Ala Gly Thr Lys Leu Glu Leu Lys Arg 225 230 235	780
gcg gcc gca gaacaaaaaac tcatctcaga agaggatctg aatggatcca Ala Ala Ala 240	829
aagacgaact ctat	843